\_\_\_\_\_\_

Sequence Listing was accepted.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: Keisha Douglas

Timestamp: [year=2008; month=1; day=17; hr=17; min=7; sec=47; ms=220; ]

\_\_\_\_\_

## Validated By CRFValidator v 1.0.3

Application No: 10532480 Version No: 2.0

Input Set:

Output Set:

**Started:** 2008-01-04 18:09:22.369

Finished: 2008-01-04 18:09:27.470

Elapsed: 0 hr(s) 0 min(s) 5 sec(s) 101 ms

Total Warnings: 69

Total Errors: 72

No. of SeqIDs Defined: 75

Actual SeqID Count: 75

| Error code | Error Description                                    |  |  |  |  |  |  |  |
|------------|--|--|--|--|--|--|--|--|
| W 213      | Artificial or Unknown found in <213> in SEQ ID (1)   |  |  |  |  |  |  |  |
| E 257      | Invalid sequence data feature in <221> in SEQ ID (1) |  |  |  |  |  |  |  |
| E 257      | Invalid sequence data feature in <221> in SEQ ID (1) |  |  |  |  |  |  |  |
| W 213      | Artificial or Unknown found in <213> in SEQ ID (2)   |  |  |  |  |  |  |  |
| E 257      | Invalid sequence data feature in <221> in SEQ ID (2) |  |  |  |  |  |  |  |
| E 257      | Invalid sequence data feature in <221> in SEQ ID (2) |  |  |  |  |  |  |  |
| W 213      | Artificial or Unknown found in <213> in SEQ ID (3)   |  |  |  |  |  |  |  |
| E 257      | Invalid sequence data feature in <221> in SEQ ID (3) |  |  |  |  |  |  |  |
| E 257      | Invalid sequence data feature in <221> in SEQ ID (3) |  |  |  |  |  |  |  |
| W 213      | Artificial or Unknown found in <213> in SEQ ID (4)   |  |  |  |  |  |  |  |
| E 257      | Invalid sequence data feature in <221> in SEQ ID (4) |  |  |  |  |  |  |  |
| E 257      | Invalid sequence data feature in <221> in SEQ ID (4) |  |  |  |  |  |  |  |
| W 213      | Artificial or Unknown found in <213> in SEQ ID (5)   |  |  |  |  |  |  |  |
| E 257      | Invalid sequence data feature in <221> in SEQ ID (5) |  |  |  |  |  |  |  |
| E 257      | Invalid sequence data feature in <221> in SEQ ID (5) |  |  |  |  |  |  |  |
| W 213      | Artificial or Unknown found in <213> in SEQ ID (6)   |  |  |  |  |  |  |  |
| E 257      | Invalid sequence data feature in <221> in SEQ ID (6) |  |  |  |  |  |  |  |
| E 257      | Invalid sequence data feature in <221> in SEQ ID (6) |  |  |  |  |  |  |  |
| W 213      | Artificial or Unknown found in <213> in SEQ ID (7)   |  |  |  |  |  |  |  |
| E 257      | Invalid sequence data feature in <221> in SEQ ID (7) |  |  |  |  |  |  |  |

## Input Set:

# Output Set:

**Started:** 2008-01-04 18:09:22.369 **Finished:** 2008-01-04 18:09:27.470

**Elapsed:** 0 hr(s) 0 min(s) 5 sec(s) 101 ms

Total Warnings: 69
Total Errors: 72
No. of SeqIDs Defined: 75

Actual SeqID Count: 75

| Error code |     | Error Description  |
|------------|-----|--|
| E          | 257 | Invalid sequence data feature in <221> in SEQ ID (7)   |
| W          | 213 | Artificial or Unknown found in <213> in SEQ ID (8)   |
| E          | 257 | Invalid sequence data feature in <221> in SEQ ID (8)   |
| E          | 257 | Invalid sequence data feature in <221> in SEQ ID (8)   |
| W          | 213 | Artificial or Unknown found in <213> in SEQ ID (9)   |
| E          | 257 | Invalid sequence data feature in <221> in SEQ ID (9)   |
| E          | 257 | Invalid sequence data feature in <221> in SEQ ID (9)   |
| W          | 213 | Artificial or Unknown found in <213> in SEQ ID (10)  |
| E          | 257 | Invalid sequence data feature in <221> in SEQ ID (10)  |
| E          | 257 | Invalid sequence data feature in <221> in SEQ ID (10) This error has occured more than 20 times, will not be displayed |
| W          | 213 | Artificial or Unknown found in <213> in SEQ ID (11)  |
| W          | 213 | Artificial or Unknown found in <213> in SEQ ID (12)  |
| W          | 213 | Artificial or Unknown found in <213> in SEQ ID (13)  |
| W          | 213 | Artificial or Unknown found in <213> in SEQ ID (14)  |
| W          | 213 | Artificial or Unknown found in <213> in SEQ ID (15)  |
| W          | 213 | Artificial or Unknown found in <213> in SEQ ID (16)  |
| W          | 213 | Artificial or Unknown found in <213> in SEQ ID (17)  |
| W          | 213 | Artificial or Unknown found in <213> in SEQ ID (18)  |
| W          | 213 | Artificial or Unknown found in <213> in SEQ ID (19)  |
| W          | 213 | Artificial or Unknown found in <213> in SEQ ID (20) This error has occured more than 20 times, will not be displayed   |
| W          | 402 | Undefined organism found in <213> in SEQ ID (66)   |

### Input Set:

# Output Set:

**Started:** 2008-01-04 18:09:22.369

Finished: 2008-01-04 18:09:27.470

**Elapsed:** 0 hr(s) 0 min(s) 5 sec(s) 101 ms

Total Warnings: 69

Total Errors: 72

No. of SeqIDs Defined: 75

Actual SeqID Count: 75

| Error code |     | Error Description |          |       |    |       |    |     |    |      |
|------------|-----|-------------------|----------|-------|----|-------|----|-----|----|------|
| W          | 402 | Undefined         | organism | found | in | <213> | in | SEQ | ID | (68) |
| W          | 402 | Undefined         | organism | found | in | <213> | in | SEQ | ID | (72) |
| W          | 402 | Undefined         | organism | found | in | <213> | in | SEQ | ID | (73) |

#### SEQUENCE LISTING

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<110> Garry, Jr., Robert F.
      McKeating, Jane A.
       Dash, Srikanta
       Coy, David H.
<120> FLAVIVIRUS FUSION INHIBITORS
<130> 12920.0014.PCUS00
<140> 10532480
<141> 2005-04-22
<150> 60/424,746
<151> 2002-11-08
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      macromolecular, t-butyloxycarbonyl, lipid, polyethyleneglycol, or
       carbohydrate
<400> 1
Tyr Gln Val Arg Asn Ser Ser Gly Leu Tyr His Val Thr Asn Asp Cys
               5
                                   10
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Pro Asn Ser Ser Ile Val Tyr Glu Ala Ala Asp Ala Ile Leu 20 25 30

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       carbohydrate
<400> 2
Cys Ser Ala Leu Tyr Trp Val Gly Asp Leu Cys Gly Ser Val Phe Leu
                5
                                    10
                                                        15
Val Gly Gln Leu Phe Thr Phe Ser Pro Arg Arg His Trp Thr Thr Gln
            2.0
                                2.5
Asp Cys
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       macromolecular, t-butyloxycarbonyl, lipid, polyethyleneglycol, or
       carbohydrate
<400> 3
Ser Pro Arg Arg His Trp Thr Thr Gln Asp Cys Asn Cys Ser Ile Tyr
               5
                                   10
Pro Gly His Ile Thr Gly His Arg Met Ala Trp Asp Met Met Asn
                               25
Trp Ser Pro Thr
       35
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       macromolecular, t-butyloxycarbonyl, lipid, polyethyleneglycol, or
       carbohydrate
<400> 4
Met Met Met Asn Trp Ser Pro Thr Ala Ala Leu Leu Arg Ile Pro Gln
               5
                                   10
                                                       1.5
Ala Ile Met Asp Met Ile Ala Gly Ala His Trp Gly Val Leu Ala Gly
```

25

30

20

35 40

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<210>
       5
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       group or one of the following groups: amido, hydrophobic,
       macromolecular, t-butyloxycarbonyl, lipid, polyethyleneglycol, or
       carbohydrate
<400> 5
Arg Val Thr Asp Pro Asp Thr Asn Thr Thr Ile Leu Thr Asn Cys Cys
                5
                                    1.0
                                                        15
Gln Arg Asn Gln Val Ile Tyr Cys Ser Pro Ser Thr Cys Leu
            20
                                25
<210> 6
<211> 35
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<222> (35)..(35)
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<400> 6

Arg Asp Phe Val Glu Gly Val Ser Gly Gly Ser Trp Val Asp Ile Val  $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$ 

Leu Glu His Gly Ser Cys Val Thr Thr Met Ala Lys Asn Lys Pro Thr 20 25 30

Leu Asp Phe

35

<210> 7

<211> 35

<212> PRT

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<400> 7

Arg Asp Phe Ile Glu Gly Ala Ser Gly Ala Thr Trp Val Asp Leu Val 1 5 10 15

Leu Glu Gly Asp Ser Cys Leu Thr Ile Met Ala Asn Asp Lys Pro Thr
20 25 30

Leu Asp Val

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<210> 8
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       group or one of the following groups: amido, hydrophobic,
       macromolecular, t-butyloxycarbonyl, lipid, polyethyleneglycol, or
       carbohydrate
<400> 8
Arg Asp Phe Ile Glu Gly Val His Gly Gly Thr Trp Val Ser Ala Thr
                5
                                   10
                                                       15
Leu Glu Gln Asp Lys Cys Val Thr Val Met Ala Pro Asp Lys Pro Ser
                                25
Leu Asp Ile
       35
<210> 9
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t-butyloxycarbonyl, lipid, polyethylene glycol, or carbohydrate

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      group or one of the following groups: amido, hydrophobic,
      macromolecular, t-butyloxycarbonyl, lipid, polyethyleneglycol, or
      carbohydrate
<400> 9
Arg Asp Phe Leu Glu Gly Val Ser Gly Ala Thr Trp Val Asp Leu Val
               5
                                    10
Leu Glu Gly Asp Ser Cys Val Thr Ile Met Ser Lys Asp Lys Pro Thr
           20
                                25
                                                    30
Ile Asp Val
        35
<210> 10
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      macromolecular, t-butyloxycarbonyl, lipid, polyethyleneglycol, or
      carbohydrate
<400> 10
Gly Gln Leu Ala Cys Lys Glu Asp Tyr Arg Tyr Ala Ile Ser Ser Thr
                5
                                    10
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Asn Glu Ile Gly Leu Leu Gly Ala Gly Gly Leu Thr Thr Thr Trp Lys
20 25 30

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Glu Tyr Asn
       35
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      group or one of the following groups: amido, hydrophobic,
      macromolecular, t-butyloxycarbonyl, lipid, polyethyleneglycol, or
      carbohydrate
<400> 11
Gly His Leu Asp Cys Lys Pro Glu Phe Ser Tyr Ala Ile Ala Lys Asp
               5
                                   10
                                                       1.5
Glu Arg Ile Gly Gln Leu Gly Ala Glu Gly Leu Thr Thr Trp Lys
           20
                                25
Glu Tyr Ser
        35
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Thr Cys Asp Ala Leu Asp Ile Gly Glu Leu Cys Gly Ala Cys Val Leu

10

15

Val Gly Asp Trp Leu Val Arg His Trp Leu Ile His Ile Asp Leu Asn 20 25 30

Glu Thr

<210> 14
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Lys Arg Phe Val Cys Lys His Ser Met Val Asp Arg Gly Trp Gly Asn 1 5 10 15

Gly Cys Gly Leu Phe Gly Lys Gly Gly Ile Val Thr Cys Ala Met Phe 20 25 30

Thr Cys

<210> 15 <211> 34 <212> PRT <213> Artificial Sequence <220> <223> Synthetic Peptide

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      macromolecular, t-butyloxycarbonyl, lipid, polyethyleneglycol, or
      carbohydrate
<400> 15
Ser Ser Tyr Val Cys Lys Gln Gly Phe Thr Asp Arg Gly Trp Gly Asn
                5
                                   10
                                                        15
Gly Cys Gly Leu Phe Gly Lys Gly Ser Ile Asp Thr Cys Ala Lys Phe
           20
                                25
Ser Cys
<210> 16
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      t-butyloxycarbonyl, lipid, polyethylene glycol, or carbohydrate
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      macromolecular, t-butyloxycarbonyl, lipid, polyethyleneglycol, or
      carbohydrate
<400> 16
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Gly Asp Asn Ala Cys Lys Arg Thr Tyr Ser Asp Arg Gly Trp Gly Asn

1 5 10 15

Gly Cys Gly Leu Phe Gly Lys Gly Ser Ile Val Ala Cys Ala Lys Phe 20 25 30

Thr Cys

<210> 17

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<400> 17

Pro Ala Phe Val Cys Arg Gln Gly Val Val Asp Arg Gly Trp Gly Asn 1 5 10 15

Gly Cys Gly Leu Phe Gly Lys Gly Ser Ile Asp Thr Cys Ala Lys Phe 20 25 30

Ala Cys

<210> 18

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<212> PRT

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- <221> MOD\_RES
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- <223> The amino-terminal amino acid residue comprises an amino group or is modified to contain one of the following groups: acetyl,